



Supplementary Fig. 2. Global transcriptome comparison of Daepung and Socheong2 at 0, 6, and 12 h after *Phytophthora sojae* isolate 2457 inoculation. (A) Correlation between RNA-seq samples. (B) Principal component analysis of transcript accumulation for 0, 6, and 12 h after *P. sojae* isolate 2457 inoculation of Daepung (DP) and Socheong2 (SC2).